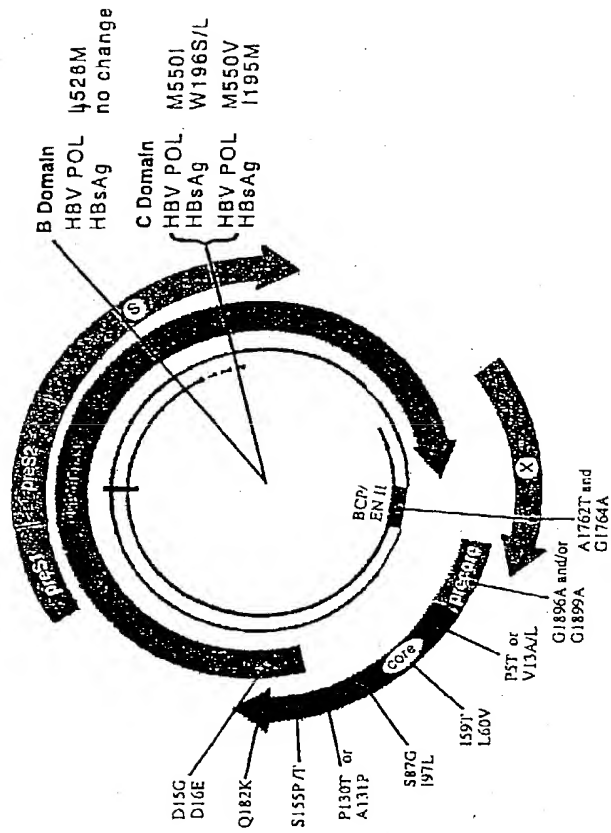


Figure 1A





**Figure 1B**

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(421) 430 440 450  
 422 438  
SNDLSWLSLD VSAAFYH<sub>1</sub>P<sub>2</sub>PL HPAAMPHELLIV GSSGLD<sub>5</sub>RYVA  
 Domain A

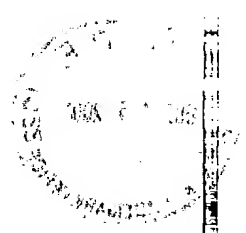
HBsAg G112R T123P Y/F134S D144E G145R  
 460 470 480 490  
 464 466 477 488 499  
 RLSST<sub>N</sub>SR<sub>N</sub>NI\*N NYHQY<sub>H</sub>G<sub>R</sub>\*\*\*D<sub>N</sub>LH D<sub>N</sub>Y<sub>S</sub>CSR<sub>D</sub>Q<sub>1</sub>LYVS LL<sub>M</sub>LLY<sub>K</sub>Q<sub>2</sub>TY<sub>F</sub>G<sub>R</sub>W

HBsAg A157D E164D F170L  
 500 510 520 530  
 512 519 523/524/526/528/530  
 KLHLY<sub>L</sub>SA<sub>H</sub>PII<sub>V</sub> LGFRKI<sub>L</sub>PMGV<sub>G</sub> GLSPFLLAQF TSAIC<sub>L</sub>SA<sub>M</sub>V<sub>T</sub>R<sub>C</sub>R  
 Domain B

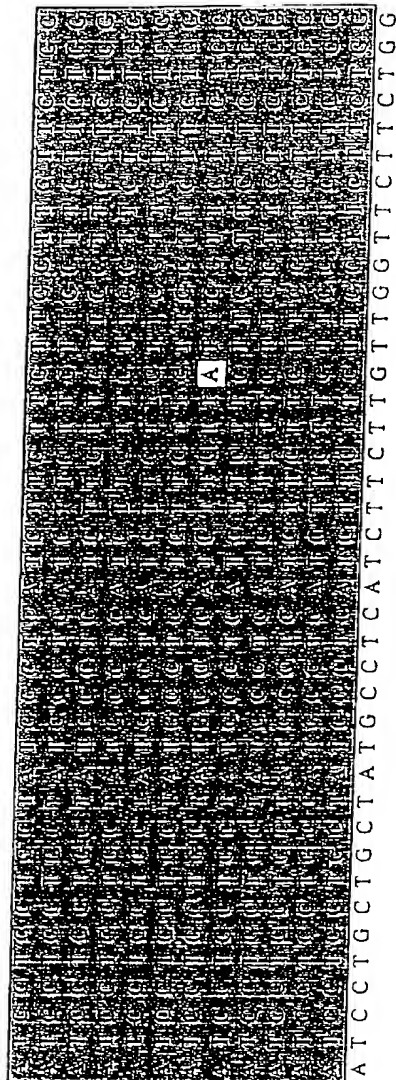
W196L W199S  
HBsAg M195I/S196W M198I S204T S210R  
 540 550 560  
 546 550 553 559 565  
 AFF<sub>P</sub>HCL<sub>V</sub>AV<sub>V</sub>FS<sub>A</sub>Y MDDV<sub>L</sub>MVLGAK<sub>R</sub>ST V<sub>G</sub>Q<sub>E</sub>HL<sub>S</sub>RES<sub>F</sub>LY<sub>F</sub>TA<sub>S</sub>A  
 Domain C

570 580 590  
 575  
 I<sub>V</sub>T<sub>C</sub>N<sub>S</sub>F<sub>V</sub>LLS<sub>D</sub>L<sub>V</sub>GI HLNPN<sub>O</sub>KTKRW GYSLNFMGYI<sub>V</sub>I G  
 Domain D Domain E

Figure 2



\*329616/HPBADR1CG  
 221499/HPBADW3  
 221500/HPBCG  
 62280/XXHEPAV  
 59439/HBVAYWE  
 59429/HBVAYWC  
 59418/HBVADW2  
 59408/HBVADRM  
 59404/HBVADR4  
 329640/HPBAYW  
 313780/HBVAYWMCG  
 229417/HPBADW1



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\*329616/HPBADR1CG  
 221499/HPBADW3  
 221500/HPBCG  
 62280/XXHEPAV  
 59439/HBVAYWE  
 59429/HBVAYWC  
 59418/HBVADW2  
 59408/HBVADRM  
 59404/HBVADR4  
 329640/HPBAYW  
 313780/HBVAYWMCG  
 229417/HPBADW1

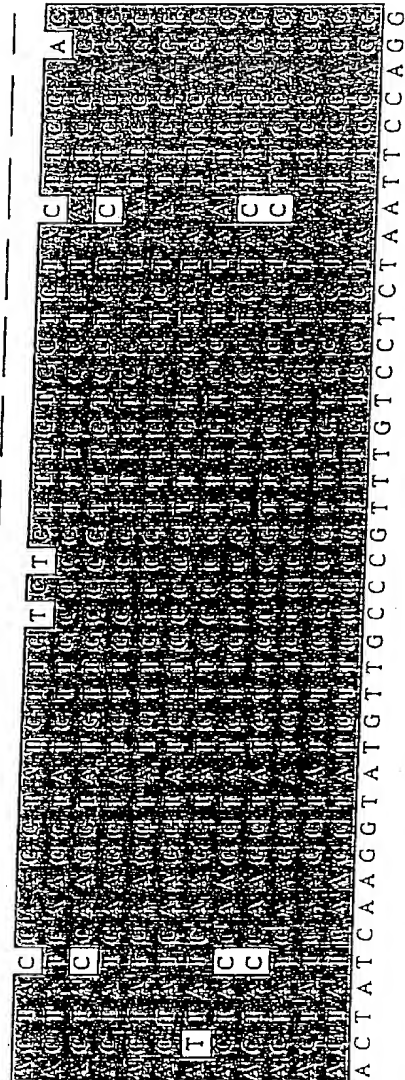
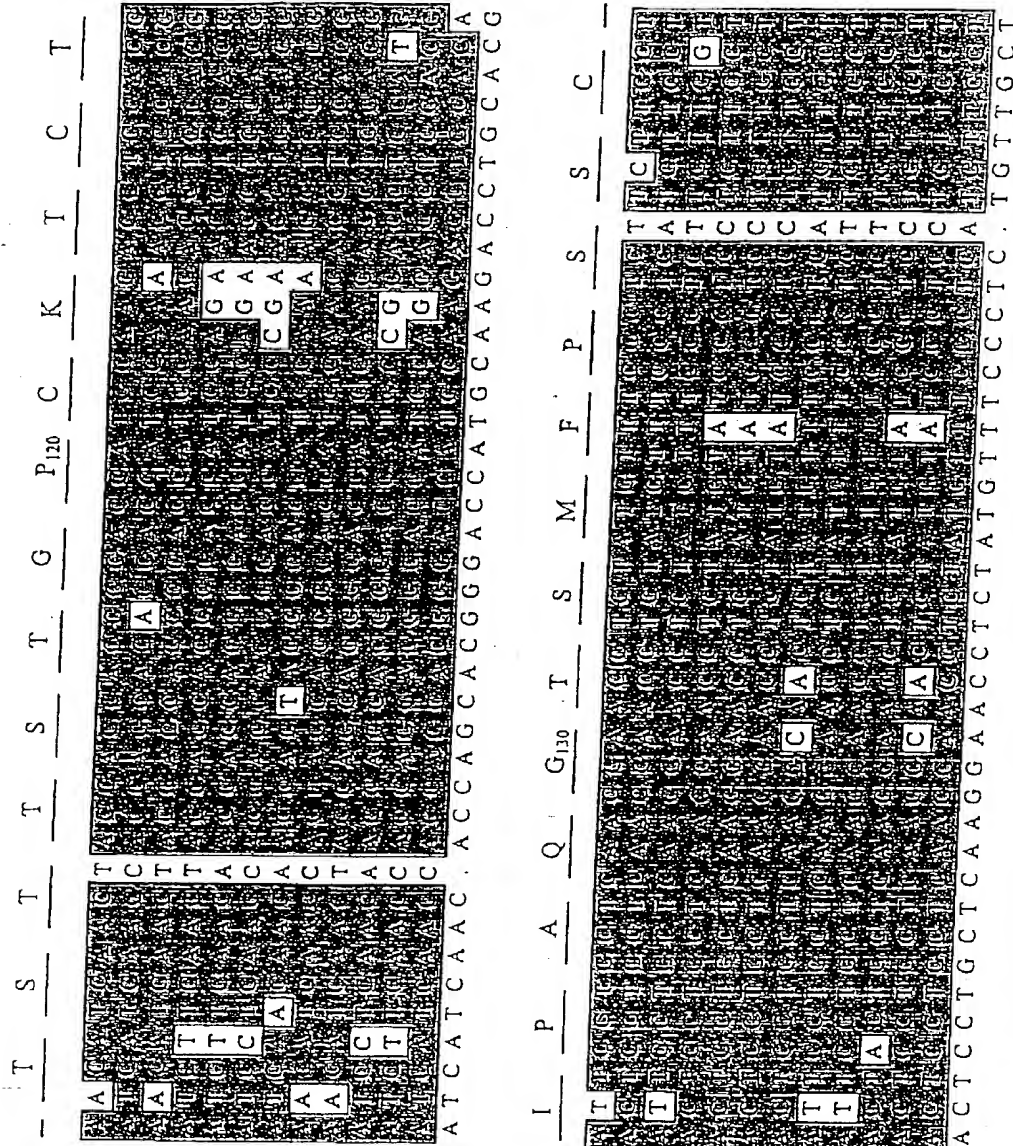


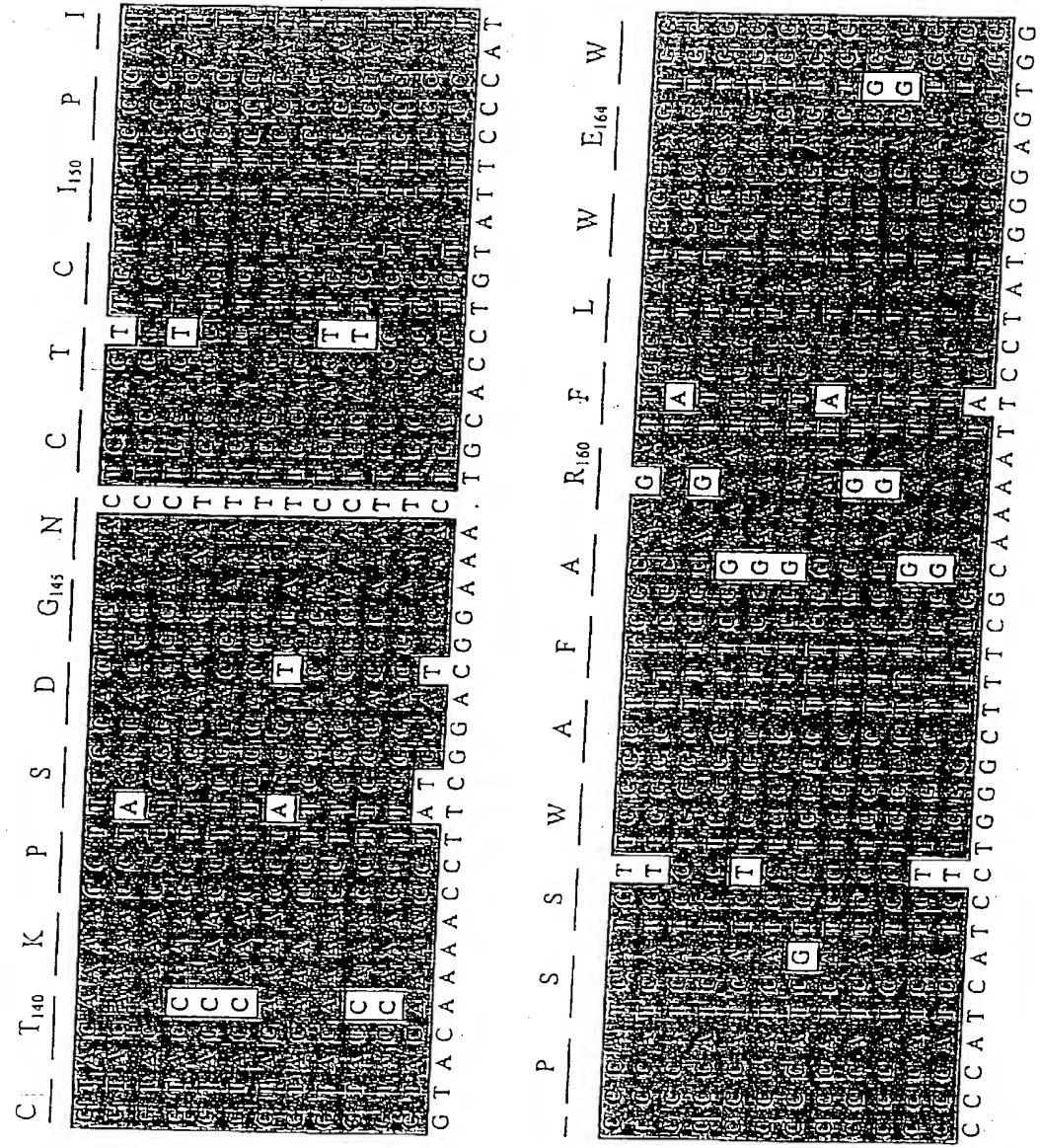
Figure 3

111



\*329616/HPBADR1CG  
 221499/HPBADW3  
 221500/HPBCG  
 62280/XXHEPAV  
 59439/HBVAYWE  
 59429/HBVAYWC  
 59418/HBVADW2  
 59408/HBVADRM  
 59404/HBVADR4  
 329640/HPBAYW  
 313780/HBVAYWMC  
 229417/HPBADW1

\*329616/HPBADR1CG  
 221499/HPBADW3  
 221500/HPBCG  
 62280/XXHEPAV  
 59439/HBVAYWE  
 59429/HBVAYWC  
 59418/HBVADW2  
 59408/HBVADRM  
 59404/HBVADR4  
 329640/HPBAYW  
 313780/HBVAYWMC  
 229417/HPBADW1



\*329616/HPBADR1CG  
221499/HPBADW3  
221500/HPBCG  
62280/XXHEPAV  
59439/HBVAYWE  
59429/HBVAYWC  
59418/HBVADW2  
59408/HBVADRM  
59404/HBVADR4  
329640/HPBAYW  
313780/HBVAYWMC  
229417/HPBADW1

\*329616/HPBADR1CG  
221499/HPBADW3  
221500/HPBCG  
62280/XXHEPAV  
59439/HBVAYWE  
59429/HBVAYWC  
59418/HBVADW2  
59408/HBVADRM  
59404/HBVADR4  
329640/HPBAYW  
313780/HBVAYWMC  
229417/HPBADW1

Figure 3 continued



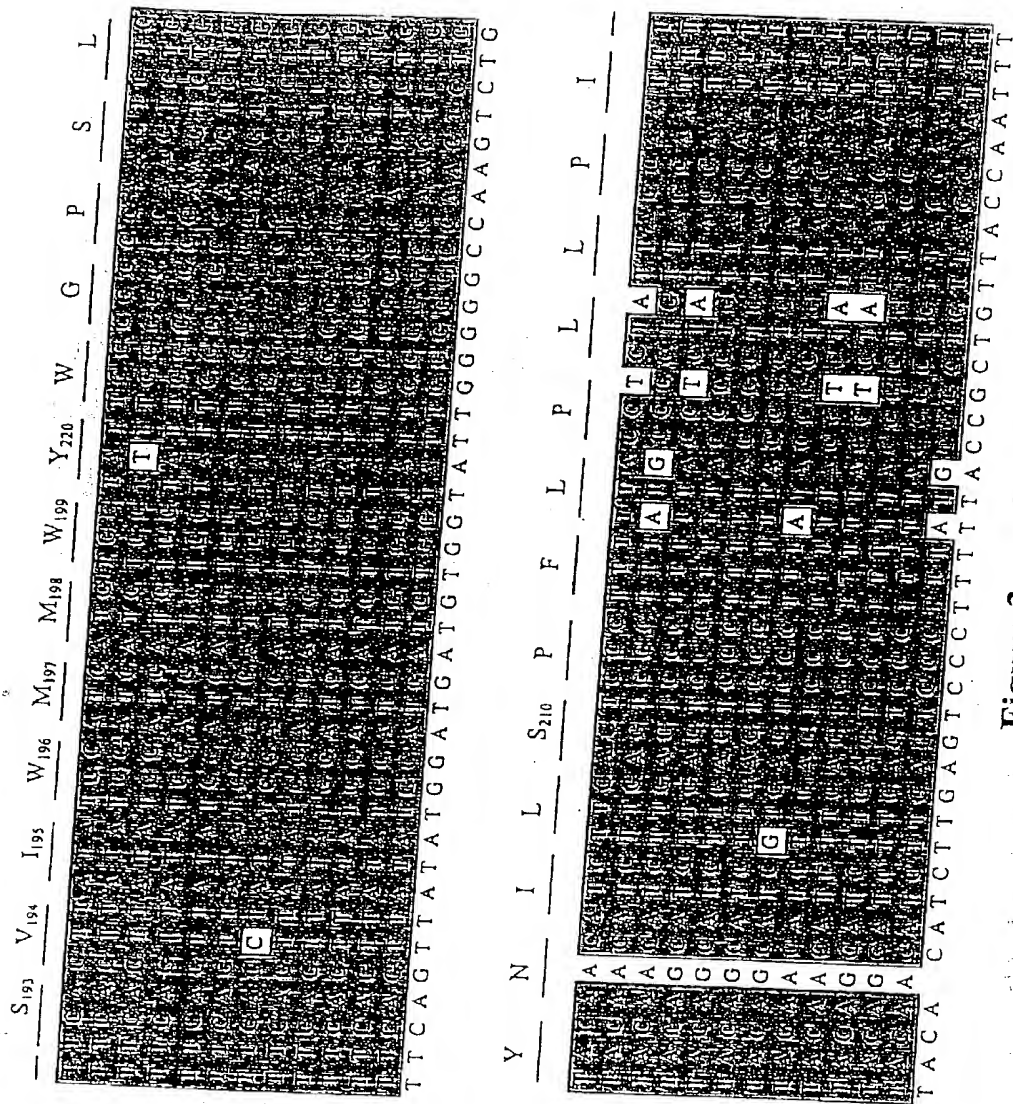
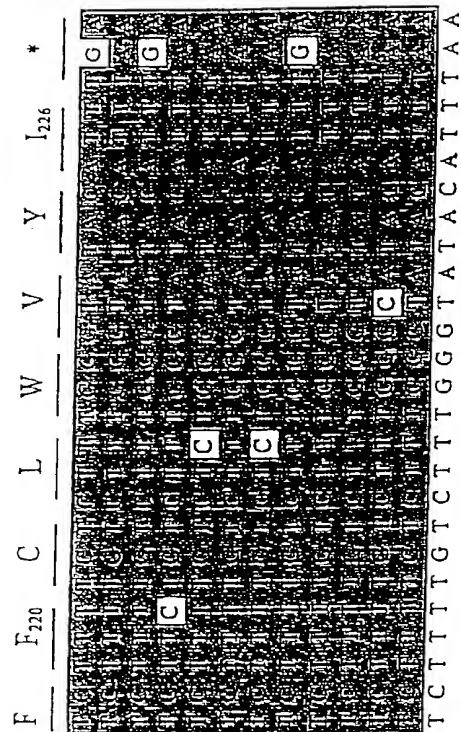


Figure 3 continued

\*329616/HPBADRICG  
 221499/HPBADW3  
 221500/HPBCG  
 62280/XXHEPAV  
 59439/HBVAYWE  
 59429/HBVAYWC  
 59418/HBVADW2  
 59408/HBVADRM  
 59404/HBVADR4  
 329640/HPBAYW  
 313780/HBVAYWMC  
 229417/HPBADW1

\*329616/HPBADRICG  
 221499/HPBADW3  
 221500/HPBCG  
 62280/XXHEPAV  
 59439/HBVAYWE  
 59429/HBVAYWC  
 59418/HBVADW2  
 59408/HBVADRM  
 59404/HBVADR4  
 329640/HPBAYW  
 313780/HBVAYWMC  
 229417/HPBADW1





\*329616/HPBADR1CG  
 221499/HPBADW3  
 221500/HPBCG  
 62280/XXHEPAV  
 59439/HBVAYWE  
 59429/HBVAYWC  
 59418/HBVADW2  
 59408/HBVADRM  
 59404/HBVADR4  
 329640/HPBAYW  
 313780/HBVAYWMCG  
 229417/HPBADW1

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Figure 3 continued

pBBHVBV1.28

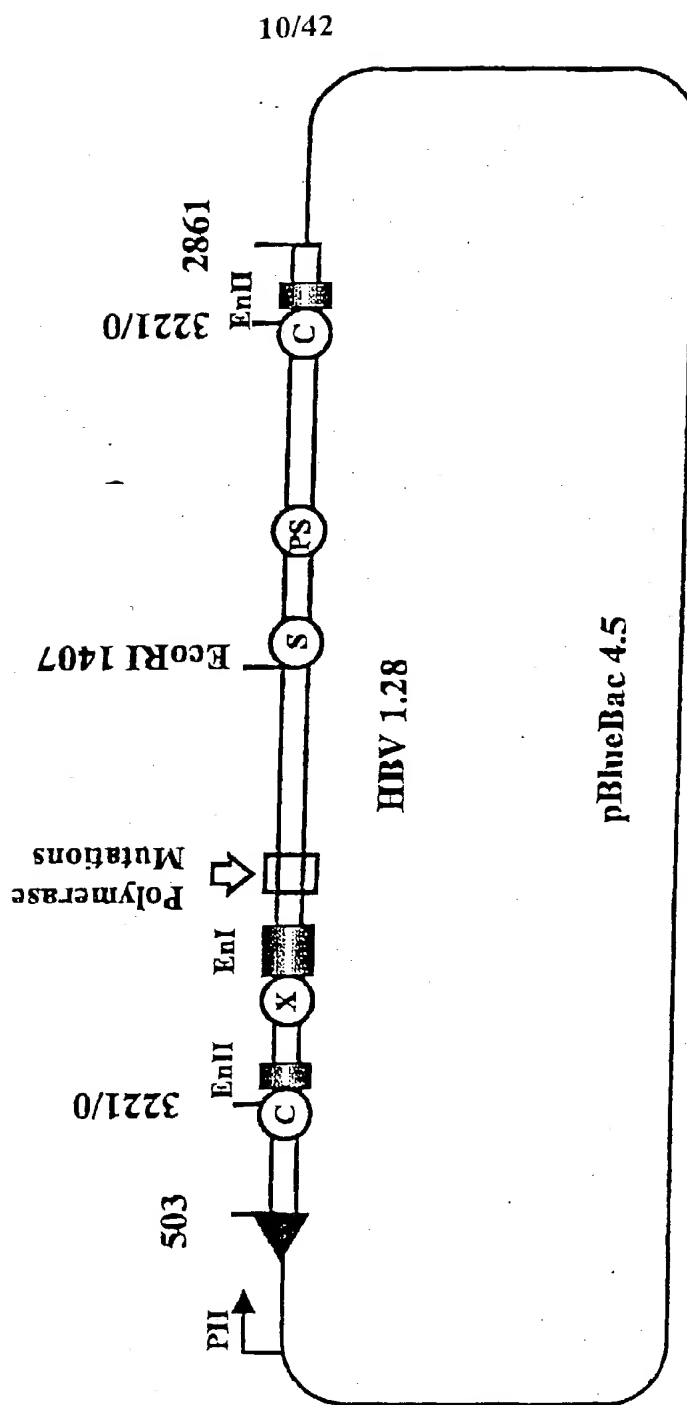


Figure 4A

pBBHBV1.5

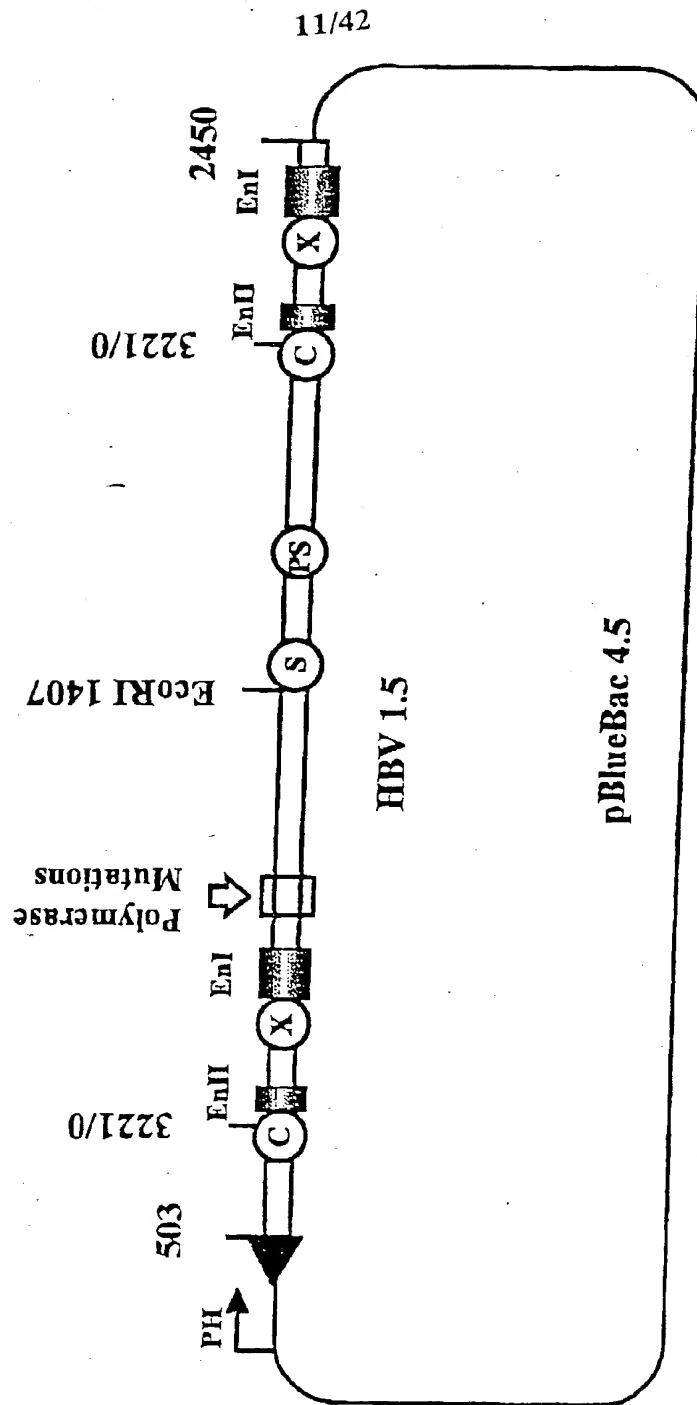


Figure 4B

12/42

Sequence Range: 1 to 4084

```

      10      20      30      40      50
GGACGACCCCTCGCGGGGCGCCTTGGGACTCTCTCGTCCCCTTCTCCGTC

      60      70      80      90     100
TGCCGTTCCAGCCGACCACGGGGCGCACCTCTCTTTACGCGGTCTCCCCG

      110     120     130     140     150
TCTGTGCCTTCTCATCTGCCGGTCCGTGTGCACTTCGCTTCACCTCTGCA

      160     170     180     190     200
CGTTGCATGGAGACCACCGTGAACGCCCATCAGATCCTGCCCAAGGTCTT

      210     220     230     240     250
ACATAAGAGGACTCTTGGACTCCCAGCAATGTCAACGACCGACCTTGAGG

      260     270     280     290     300
CCTACTTCAAAGACTGTGTGTTTAAGGACTGGGAGGAGCTGGGGGAGGAG

      310     320     330     340     350
ATTAGGTAAAGGTCTTTGTATTAGGAGGCTGTAGGCATAAATTGGTCTG

      360     370     380     390     400
CGCACCAGCACCATGCAACTTTTTACCTCTGCCTAATCATCTCTTGTA

      410     420     430     440     450
ATGTCCCCTGTTCAAGCCTCCAAGCTGTGCCTTGGGTGGCTTTGGGGCA

      460     470     480     490     500
TGGACATTGACCCTTATAAAGAATTTGGAGCTACTGTGGAGTTACTCTCG

      510     520     530     540     550
TTTTTGCTTCTGACTTCTTTCCTTCCGTCAGAGATCTCCTAGACACCGC

      560     570     580     590     600
CTCAGCTCTGTATCGAGAAGCCTTAGAGTCTCCTGAGCATTGCTCACCTC

      610     620     630     640     650
ACCATACTGCACTCAGGCAAGCCATTCTCTGCTGGGGGGAATTGATGACT

      660     670     680     690     700
CTAGCTACCTGGGTGGGTAATAATTTGGAAGATCCAGCATCCAGGGATCT

```

Figure 5A

710 720 730 740 750  
AGTAGTCAATTATGTTAATACTAACATGGGTTTAAAGATCAGGCAACTAT

760 770 780 790 800  
TGTGGTTTTCATATATCTTGCCTTACTTTTGAAGAGAGACTGTACTTGAA

810 820 830 840 850  
TATTTGGTCTCTTTTCGGAGTGTGGATTTCGCACTCCTCCAGCCTATAGACC

860 870 880 890 900  
ACCAAATGCCCTATCTTATCAACACTTCCGGAACTACTGTTGTTAGAC

910 920 930 940 950  
GACGGGACCGAGGCAGGTCCCCTAGAAGAAGAACTCCCTCGCCTCGCAGA

960 970 980 990 1000  
CGCAGATCTCAATCGCCGCGTCGCAGAAGATCTCAATCTCGGAATCTCA

1010 1020 1030 1040 1050  
ATGTTAGTATTCCTTGGACTCATAAGGTGGGAACTTTACGGGGCTTTAT

1060 1070 1080 1090 1100  
TCCTCTACAGTACCTATCTTTAATCCTGAATGGCAAACCTCCTTCCTTTCC

1110 1120 1130 1140 1150  
TAAGATTCAATTTACAAGAGGACATTATTAATAGGTGTCAACAATTTGTGG

1160 1170 1180 1190 1200  
GCCCTCTCACTGTAAATGAAAAGAGAAGATTGAAATTAATTATGCCTTGCT

1210 1220 1230 1240 1250  
AGATTCTATCCTACCCACACTAAATATTTGCCCTTAGACAAAGGAATTAA

1260 1270 1280 1290 1300  
ACCTTATTATCCAGATCAGGTAGTTAATCATTACTTCCAAACCAGACATT

1310 1320 1330 1340 1350  
ATTACATACTCTTTGGAAGGCTGGTATTCTATATAAGAGGGAAACCACA

1360 1370 1380 1390 1400  
CGTAGCGCATCATTTTGCGGGTCAACCATATTCTTGGAACAAGAGCTACA

1410 1420 1430 1440 1450  
GCATGGGAGGTTGGTCATCAAAACCTCGCAAAGGCATGGGGACGAATCTT

Figure 5A continued

14/42

1460 1470 1480 1490 1500  
TCTGTTCCCAACCCTCTGGGATTCTTTCCCGATCATCAGTTGGACCCCTGC

1510 1520 1530 1540 1550  
ATTTCGGAGCCAACCTCAAACAATCCAGATTGGGACTTCAACCCCATCAAGG

1560 1570 1580 1590 1600  
ACCACTGGCCAGCAGCCAACCAGGTAGGAGTGGGAGCATTCTGGGCCAGGG

1610 1620 1630 1640 1650  
CTCACCCCTCCACACGGCGGTATTTGGGGTGGAGCCCTCAGGCTCAGGG

1660 1670 1680 1690 1700  
CATATTGACCACAGTGTCAACAATTCCTCCTCCTGCCTCCACCAATCGGC

1710 1720 1730 1740 1750  
AGTCAGGAAGGCAGCCTACTCCCATCTCTCCACCTCTAAGAGACAGTCAT

1760 1770 1780 1790 1800  
CCTCAGGCCATGCAGTGGAAATTCCTGCTTCCACCAAGCTCTGCAGGA

1810 1820 1830 1840 1850  
TCCCAGAGTCAGGGGTCTGTATCTTCTGCTGGTGGCTCCAGTTCAGGAA

1860 1870 1880 1890 1900  
CAGTAAACCCTGCTCCGAATATTGCCTCTCACATCTCGTCAATCTCCGCG

1910 1920 1930 1940 1950  
AGGACTGGGGACCCTGTGACGAACATGGAGAACATCACATCAGGATTCTCT

1960 1970 1980 1990 2000  
AGGACCCCTGCTCGTGTTACAGGCGGGGTTTTCTTGTTGACAAGAATCC

2010 2020 2030 2040 2050  
TCACAATACCGCAGAGTCTAGACTCGTGGTGGACTTCTCTCAATTTTCTA

2060 2070 2080 2090 2100  
GGGGGATCTCCCGTGTGTCTTGCCAAAATTCCGAGTCCCCAACCTCCAA

2110 2120 2130 2140 2150  
TCACTCACCAACCTCCTGTCTCCAATTTGTCCTGGTTATCGCTGGATGT

2160 2170 2180 2190 2200  
GTCTGCGGCGTTTTATCATATTCTCTTCATCCTGCTGCTATGCCTCATC

Figure 5A continued

15/42

2210 2220 2230 2240 2250  
TTCTTATTGGTTCTTCTGGATTATCAAGGTATGTTGCCCGTTTGTCTCT

2260 2270 2280 2290 2300  
AATTCCAGGATCAACAACAACCAAGTACGGGACCATGCAAAACCTGCACGA

2310 2320 2330 2340 2350  
CTCCTGCTCAAGGCAACTCTATGTTTCCCTCATGTTGCTGTACAAAACCT

2360 2370 2380 2390 2400  
ACGGATGGAAATTGCACCTGTATCCCATCCCATCGTCCTGGGCTTTTCGC

2410 2420 2430 2440 2450  
AAAATACCTATGGGAGTGGGCCTCAGTCCGTTTCTCTTGGCTCAGTTTAC

2460 2470 2480 2490 2500  
TAGTGCCATTTGTTCAAGTGGTTCGTAGGGCTTTCCCCCACTGTTTGGCTT

2510 2520 2530 2540 2550  
TCAGCTATATGGATGATGTGCTATTGGGGGCCAAGTCTGTACAGCATCGT

2560 2570 2580 2590 2600  
GAGTCCCTTTATACCGCTGTTACCAATTTTCTTTTGTCTCTGGGTATACA

2610 2620 2630 2640 2650  
TTTAAACCCTAACAACCAACCAAGATGGGGTTATTCCCTAAACTTCATGG

2660 2670 2680 2690 2700  
GCTACATAATTGGAAGTTGGGGAACCTTGCCACAGGATCATATTGTACAA

2710 2720 2730 2740 2750  
AAGATCAAACACTGTTTTAGAAAACCTCCTGTTAACAGGCCTATTGATTG

2760 2770 2780 2790 2800  
GAAAGTATGTCAAAGAATTGTGGGTCTTTTGGGCTTTGCTGCTCCATTTA

2810 2820 2830 2840 2850  
CACAATGTGGATATCCTGCCTTAATGCCTTTGTATGCATGTATACAAGCT

2860 2870 2880 2890 2900  
AAACAGGCTTTCACTTTCTCGCCAACCTTACAAGGCCTTTCTAAGTAAACA

2910 2920 2930 2940 2950  
GTACATGAACCTTTACCCCGTTGCTCGGCAACGGCCTGGTCTGTGCCAAG

Figure 5A continued

16/42

2960 2970 2980 2990 3000  
TGTTTGCTGACGCAACCCCCACTGGCTGGGGCTTGGCCATAGGCCATCAG  
3010 3020 3030 3040 3050  
CGCATGCGTGGAACCTTTGTGGCTCCTCTGCCGATCCATACTGCGGAACT  
3060 3070 3080 3090 3100  
CCTAGCCGCTTGTTTTGCTCGCAGCCGGTCTGGAGCAAAGCTCATCGGAA  
3110 3120 3130 3140 3150  
CTGACAATTCTGTCGTCCTCTCGCGGAAATATACATCGTTTCCATGGCTG  
3160 3170 3180 3190 3200  
CTAGGCTGTACTGCCAACTGGATCCTTCGCGGGACGTCCTTTGTTTACGT  
3210 3220 3230 3240 3250  
CCCGTCGGCGCTGAATCCCGCGGACGACCCCTCGCGGGGCCGCTTGGGAC  
3260 3270 3280 3290 3300  
TCTCTCGTCCCCTTCTCCGTCTGCCGTTCCAGCCGACCACGGGGCGCACC  
3310 3320 3330 3340 3350  
TCTCTTTACGCGGTCTCCCCGTCTGTGCCTTCTCATCTGCCGGTCCGTGT  
3360 3370 3380 3390 3400  
GCACTTCGCTTCACCTCTGCACGTTGCATGGAGACCACCGTGAACGCCCA  
3410 3420 3430 3440 3450  
TCAGATCCTGCCCAAGGTCTTACATAAGAGGACTCTTGGACTCCCAGCAA  
3460 3470 3480 3490 3500  
TGTCAACGACCGACCTTGAGGCCTACTTCAAAGACTGTGTGTTTAAGGAC  
3510 3520 3530 3540 3550  
TGGGAGGAGCTGGGGGAGGAGATTAGGTTAAAGGTCTTTGTATTAGGAGG  
3560 3570 3580 3590 3600  
CTGTAGGCATAAATTGGTCTGCGCACCAGCACCATGCAACTTTTTTACCT  
3610 3620 3630 3640 3650  
CTGCCTAATCATCTCTTGTACATGTCCCCTGTTCAAGCCTCCAAGCTGT  
3660 3670 3680 3690 3700  
GCCTTGGGTGGCTTTGGGGCATGGACATTGACCCTTATAAAGAATTTGGA

Figure 5A continued



17/42

3710 3720 3730 3740 3750  
GCTACTGTGGAGTTACTCTCGTTTTTGCCTTCTGACTTCTTTCCTTCCGT

3760 3770 3780 3790 3800  
CAGAGATCTCCTAGACACCGCCTCAGCTCTGTATCGAGAAGCCTTAGAGT

3810 3820 3830 3840 3850  
CTCCTGAGCATTGCTCACCTCACCATACTGCACTCAGGCAAGCCATTCTC

3860 3870 3880 3890 3900  
TGCTGGGGGGAATTGATGACTCTAGCTACCTGGGTGGGTAATAATTTGGA

3910 3920 3930 3940 3950  
AGATCCAGCATCCAGGGATCTAGTAGTCAATTATGTTAATACTAACATGG

3960 3970 3980 3990 4000  
GTTTAAAGATCAGGCAACTATTGTGGTTTCATATATCTTGCCTTACTTTT

4010 4020 4030 4040 4050  
GGAAGAGAGACTGTACTTGAATATTTGGTCTCTTTCGGAGTGTGGATTCTG

4060 4070 4080  
CACTCCTCCAGCCTATAGACCACCAAATGCCCT

Figure 5A continued

18/42

Sequence Range: 1 to 4496

```
      10      20      30      40      50
GATATCCTGCCTTAATGCCTTTGTATGCATGTATACAAGCTAAACAGGCT

      60      70      80      90     100
TTCACTTTCTCGCCAACCTACAAGGCCTTTCTAAGTAAACAGTACATGAA

      110     120     130     140     150
CCTTTACCCCGTTGCTCGGCAACGGCCTGGTCTGTGCCAAGTGTGCTG

      160     170     180     190     200
ACGCAACCCCACTGGCTGGGGCTTGGCCATAGGCCATCAGCGCATGCGT

      210     220     230     240     250
GGAACCTTTGTGGCTCCTCTGCCGATCCATACTGCGGAACCTCTAGCCGC

      260     270     280     290     300
TTGTTTGTCTCGCAGCCGGTCTGGAGCAAAGCTCATCGGAACCTGACAATT

      310     320     330     340     350
CTGTCTGTCCTCTCGCGGAAATATACATCGTTTCCATGGCTGCTAGGCTGT

      360     370     380     390     400
ACTGCCAACTGGATCCTTCGCGGGACGTCCTTTGTTTACGTCCCGTCGGC

      410     420     430     440     450
GCTGAATCCCGCGGACGACCCCTCGCGGGGCGCGCTTGGGACTCTCTCGTC

      460     470     480     490     500
CCCTTCTCCGTCTGCCGTTCCAGCCGACCACGGGGCGCACCTCTCTTTAC

      510     520     530     540     550
GCGGTCTCCCGTCTGTGCCTTCTCATCTGCCGGTCCGTGTGCACTTCGC

      560     570     580     590     600
TTCACCTCTGCACGTTGCATGGAGACCACCGTGAACGCCCATCAGATCCT

      610     620     630     640     650
GCCCAAGGTCTTACATAAGAGGACTCTTGGACTCCCAGCAATGTCAACGA

      660     670     680     690     700
CCGACCTTGAGGCCTACTTCAAAGACTGTGTGTTTAAGGACTGGGAGGAG
```

Figure 5B

19/42

710 720 730 740 750  
CTGGGGGAGGAGATTAGGTTAAAGGTCTTTGTATTAGGAGGCTGTAGGCA  
760 770 780 790 800  
TAAATTGGTCTGCGCACCAGCAGCATGCAACTTTTTACCTCTGCCTAAT  
810 820 830 840 850  
CATCTCTTGTACATGTCCCACTGTTCAAGCCTCCAAGCTGTGCCTTGGGT  
860 870 880 890 900  
GGCTTTGGGGCATGGACATTGACCCTTATAAAGAATTGGAGCTACTGTG  
910 920 930 940 950  
GAGTTACTCTCGTTTTTGCCTTCTGACTTCTTTCCTCCGTCAGAGATCT  
960 970 980 990 1000  
CCTAGACACCGCCTCAGCTCTGTATCGAGAAGCCTTAGAGTCTCCTGAGC  
1010 1020 1030 1040 1050  
ATTGCTCACCTCACCATACTGCACTCAGGCAAGCCATTCTCTGCTGGGGG  
1060 1070 1080 1090 1100  
GAATTGATGACTCTAGCTACCTGGGTGGGTAATAATTTGGAAGATCCAGC  
1110 1120 1130 1140 1150  
ATCCAGGGATCTAGTAGTCAATTATGTTAATACTAACATGGGTTTAAAGA  
1160 1170 1180 1190 1200  
TCAGGCAACTATTGTGGTTTTCATATATCTTGCCTTACTTTTGGAAGAGAG  
1210 1220 1230 1240 1250  
ACTGTACTTGAATATTTGGTCTCTTTCGGAGTGTGGATTTCGCACTCCTCC  
1260 1270 1280 1290 1300  
AGCCTATAGACCACCAAATGCCCTATCTTATCAACACTTCCGGAAACTA  
1310 1320 1330 1340 1350  
CTGTTGTTAGACGACGGGACCGAGGCAGGTCCCCTAGAAGAAGAACTCCC  
1360 1370 1380 1390 1400  
TCGCCTCGCAGACGCAGATCTCAATCGCCGCGTCGCAGAAGATCTCAATC  
1410 1420 1430 1440 1450  
TCGGGAATCTCAATGTTAGTATTCCTTGGACTCATAAGGTGGGAACTTT

Figure 5B continued

20/42

1460 1470 1480 1490 1500  
ACGGGGCTTTATTCTCTACAGTACCTATCTTTAATCCTGAATGGCAAAC

1510 1520 1530 1540 1550  
TCCTTCCTTTTCTAAGATTCATTTACAAGAGGACATTATTAATAGGTGTC

1560 1570 1580 1590 1600  
AACAATTTGTGGGCCCTCTCACTGTAAATGAAAAGAGAAGATTGAAATTA

1610 1620 1630 1640 1650  
ATTATGCCTGCTAGATTCTATCCTACCCACACTAAATATTTGCCCTTAGA

1660 1670 1680 1690 1700  
CAAAGGAATTAAACCTTATTATCCAGATCAGGTAGTTAATCATTACTTCC

1710 1720 1730 1740 1750  
AAACCAGACATTATTTACATACTCTTTGGAAGGCTGGTATTCTATATAAG

1760 1770 1780 1790 1800  
AGGGAAACCACACGTAGCGCATCTTTGCGGGTCACCATATTCTTGGGA

1810 1820 1830 1840 1850  
ACAAGAGCTACAGCATGGGAGGTTGGTCATCAAAACCTCGCAAAGGCATG

1860 1870 1880 1890 1900  
GGGACGAATCTTTCTGTTCCTCAACCTCTGGGATTCTTTCCCGATCATCA

1910 1920 1930 1940 1950  
GTTGGACCCTGCATTTCGGAGCCAACTCAAACAATCCAGATTGGGACTTCA

1960 1970 1980 1990 2000  
ACCCCATCAAGGACCACTGGCCAGCAGCCAACCAGGTAGGAGTGGGAGCA

2010 2020 2030 2040 2050  
TTCGGGCCAGGGCTCACCCCTCCACACGGCGGTATTTTGGGGTGGAGCCC

2060 2070 2080 2090 2100  
TCAGGCTCAGGGCATATTGACCACAGTGTCAACAATTCTCCTCCTGCCT

2110 2120 2130 2140 2150  
CCACCAATCGGCAGTCAGGAAGGCAGCCTACTCCCATCTCTCCACCTCTA

2160 2170 2180 2190 2200  
AGAGACAGTCATCCTCAGGCCATGCAGTGAATTCCACTGCCTTCCACCA

Figure 5B continued

21/42

2210 2220 2230 2240 2250  
AGCTCTGCAGGATCCCAGAGTCAGGGGTCTGTATCTTCCTGCTGGTGGCT

2260 2270 2280 2290 2300  
CCAGTTCAGGAACAGTAAACCCTGCTCCGAATATTGCCTCTCACATCTCG

2310 2320 2330 2340 2350  
TCAATCTCCGCGAGGACTGGGGACCCTGTGACGAACATGGAGAACATCAC

2360 2370 2380 2390 2400  
ATCAGGATTCTTAGGACCCCTGCTCGTGTACAGGCGGGGTTTTCTTGT

2410 2420 2430 2440 2450  
TGACAAGAATCCTCACAATACCGCAGAGTCTAGACTCGTGGTGGACTTCT

2460 2470 2480 2490 2500  
CTCAATTTTCTAGGGGGATCTCCCGTGTGTCTTGGCCAAAATTGCGAGTC

2510 2520 2530 2540 2550  
CCCAACCTCCAATCACTCACCAACCTCCTGTCTCTCCAATTTGTCCTGGTT

2560 2570 2580 2590 2600  
ATCGCTGGATGTGTCTGCGGCGTTTTATCATATTCTCTTCATCCTGCTG

2610 2620 2630 2640 2650  
CTATGCCTCATCTTCTTATTGGTTCTTCTGGATTATCAAGGTATGTTGCC

2660 2670 2680 2690 2700  
CGTTTGTCTCTAATTCCAGGATCAACAACAACCAGTACGGGACCATGCA

2710 2720 2730 2740 2750  
AAACCTGCACGACTCCTGCTCAAGGCAACTCTATGTTTCCCTCATGTTGC

2760 2770 2780 2790 2800  
TGTACAAAACCTACGGATGGAAATTGCACCTGTATTCCCATCCCATCGTC

2810 2820 2830 2840 2850  
CTGGGCTTTTCGAAAATACCTATGGGAGTGGGCCTCAGTCCGTTTCTCTT

2860 2870 2880 2890 2900  
GGCTCAGTTTACTAGTGCCATTTGTTTCAGTGGTTTCGTAGGGCTTTCCCCC

2910 2920 2930 2940 2950  
ACTGTTTGGCTTTTCAGCTATATGGATGATGTGGTATTGGGGGCCAAGTCT

Figure 5B continued

22/42

2960 2970 2980 2990 3000  
GTACAGCATCGTGAGTCCCTTTATACCGCTGTTACCAATTTTCTTTTGTC

3010 3020 3030 3040 3050  
TCTGGGTATACATTTAAACCCTAACAAAACAAAAGATGGGGTTATTCCC

3060 3070 3080 3090 3100  
TAAACTTCATGGGCTACATAATTGGAAGTTGGGGAACTTTGCCACAGGAT

3110 3120 3130 3140 3150  
CATATTGTACAAAAGATCAAACACTGTTTGTAGAAAACCTCCTGTTAACAG

3160 3170 3180 3190 3200  
GCCTATTGATTGGAAAGTATGTCAAAGAATTGTGGGTCTTTTGGGCTTTG

3210 3220 3230 3240 3250  
CTGCTCCATTTACACAATGTGGATATCCTGCCTTAATGCCTTTGTATGCA

3260 3270 3280 3290 3300  
TGTATACAAGCTAAACAGGCTTTCACTTTCTCGCCAACCTTACAAGGCCTT

3310 3320 3330 3340 3350  
TCTAAGTAAACAGTACATGAACCTTTACCCCGTTGCTCGGCAACGGCCTG

3360 3370 3380 3390 3400  
GTCTGTGCCAAGTGTTTGCTGACGCAACCCCCACTGGCTGGGGCTTGGCC

3410 3420 3430 3440 3450  
ATAGGCCATCAGCGCATGCGTGGAACCTTTGTGGCTCCTCTGCCGATCCA

3460 3470 3480 3490 3500  
TACTGCGGAACCTCCTAGCCGCTTGTGCTCGCAGCCGGTCTGGAGCAA

3510 3520 3530 3540 3550  
AGCTCATCGGAACCTGACAATTCTGTCTCCTCTCGCGGAAATATACATCG

3560 3570 3580 3590 3600  
TTTCCATGGCTGCTAGGCTGTACTGCCAACTGCATCCTTCGCGGGACGTC

3610 3620 3630 3640 3650  
CTTTGTTTACGTCCCGTCGGCGCTGAATCCCGCGGACGACCCCTCGCGGG

3660 3670 3680 3690 3700  
GCCGCTTGGGACTCTCTCGTCCCCCTTCTCCGTCTGCCGTTCCAGCCGACC

Figure 5B continued

3710 3720 3730 3740 3750  
ACGGGGCGCACCTCTCTTTACGCGGTCTCCCCGTCTGTGCCTTCTCATCT

3760 3770 3780 3790 3800  
GCCGGTCCGTGTGCACTTCGCTTCACCTCTGCACGTTGCATGGAGACCAC

3810 3820 3830 3840 3850  
CGTGAACGCCCCATCAGATCCTGCCCAAGGTCTTACATAAGAGGACTCTTG

3860 3870 3880 3890 3900  
GACTCCCAGCAATGTCAACGACCGACCTTGAGGCCTACTTCAAAGACTGT

3910 3920 3930 3940 3950  
GTGTTTAAGGACTGGGAGGAGCTGGGGGAGGAGATTAGGTTAAAGGTCTT

3960 3970 3980 3990 4000  
TGTATTAGGAGGCTGTAGGCATAAATTGGTCTGCGCACCAGCACCATGCA

4010 4020 4030 4040 4050  
ACTTTTTACCTCTGCCTAATCATCTCTTGTACATGTCCCACTGTTCAAG

4060 4070 4080 4090 4100  
CCTCCAAGCTGTGCCTTGGGTGGCTTTGGGGCATGGACATTGACCCTTAT

4110 4120 4130 4140 4150  
AAAGAATTTGGAGCTACTGTGGAGTTACTCTCGTTTTTGCCTTCTGACTT

4160 4170 4180 4190 4200  
CTTTCCTTCCGTCAGAGATCTCCTAGACACCGCCTCAGCTCTGTATCGAG

4210 4220 4230 4240 4250  
AAGCCTTAGAGTCTCCTGAGCATTGCTCACCTCACCATACTGCACTCAGG

4260 4270 4280 4290 4300  
CAAGCCATTCTCTGCTGGGGCGAATTGATGACTCTAGCTACCTGGGTGGG

4310 4320 4330 4340 4350  
TAATAATTTGGAAGATCCAGCATCCAGGGATCTAGTAGTCAATTATGTTA

4360 4370 4380 4390 4400  
ATACTAACATGGGTTTAAAGATCAGGCAACTATTGTGGTTTCATATATCT

4410 4420 4430 4440 4450  
TGCCTTACTTTTGAAGAGAGACTGTACTTGAATATTTGGTCTCTTTTCGG

4460 4470 4480 4490  
AGTGTGGATTGCACTCCTCCAGCCTATAGACCACCAAATGCCCT

Figure 5B continued

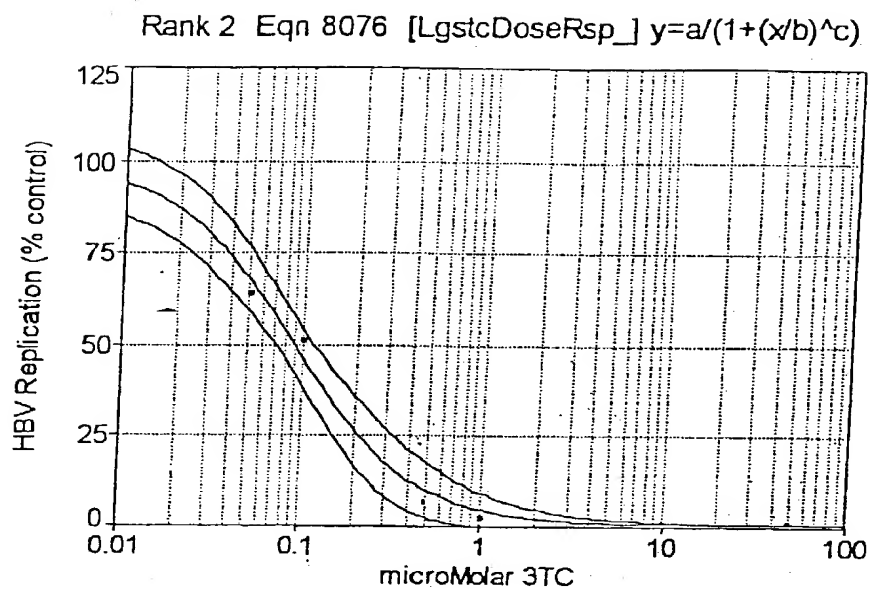


Figure 6A

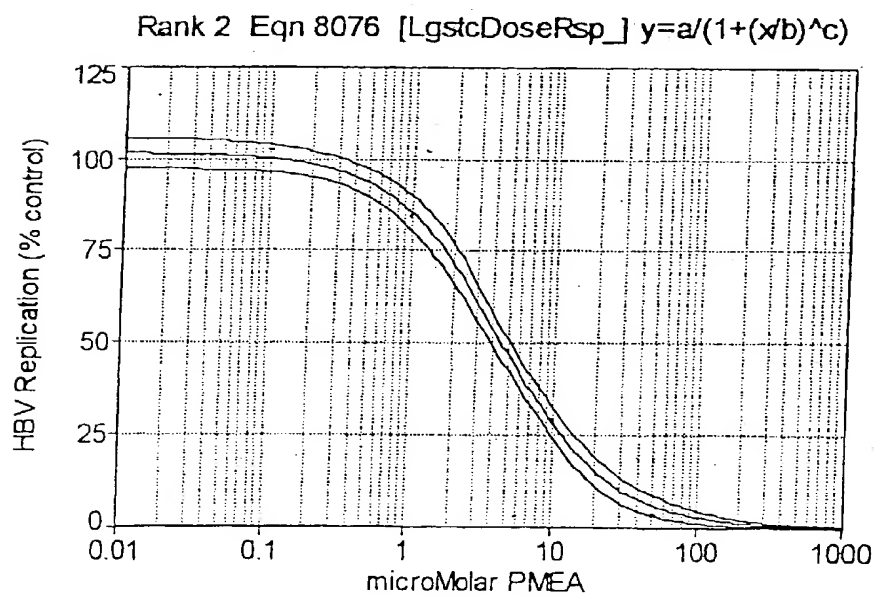


Figure 6B



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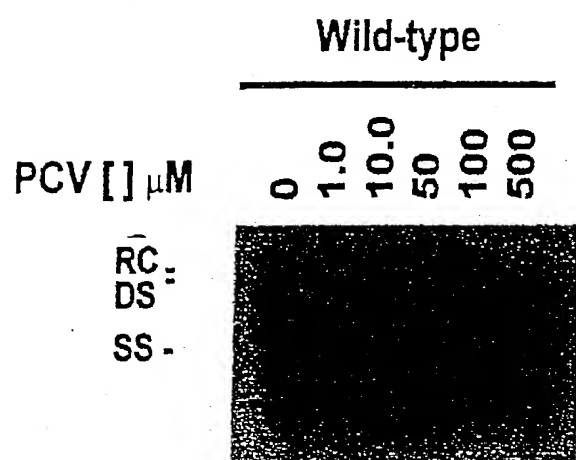


Figure 6C

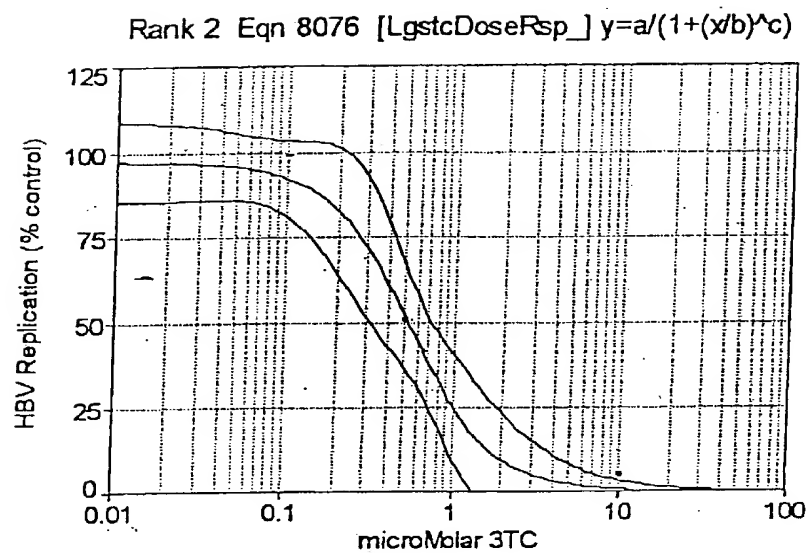


Figure 7A

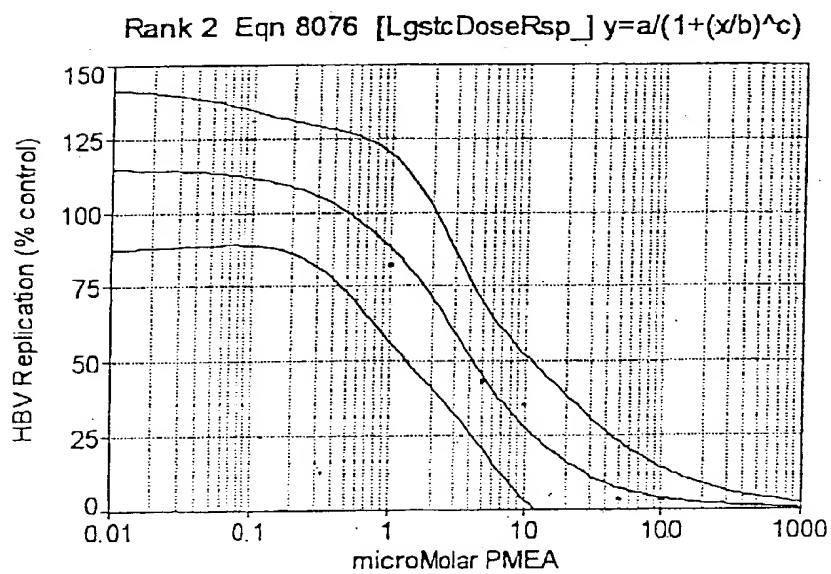


Figure 7B

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Rank 45 Eqn 19  $y=a+b\ln x/x^2$

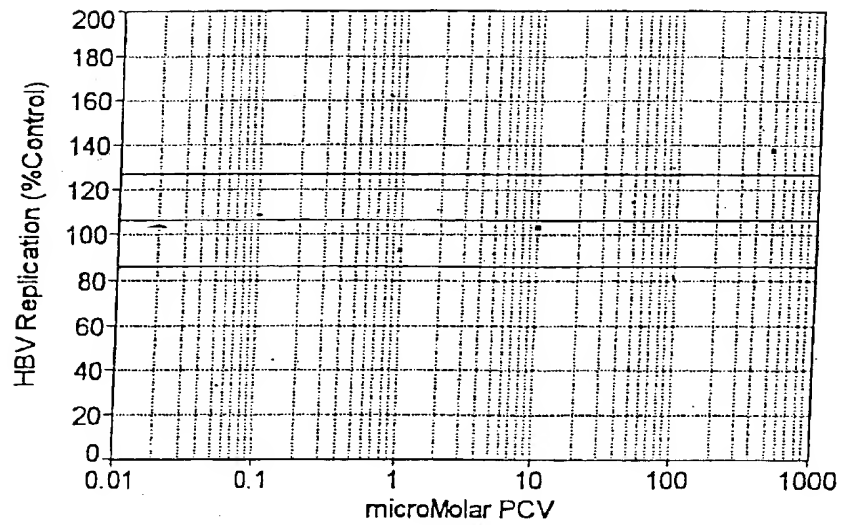


Figure 7C

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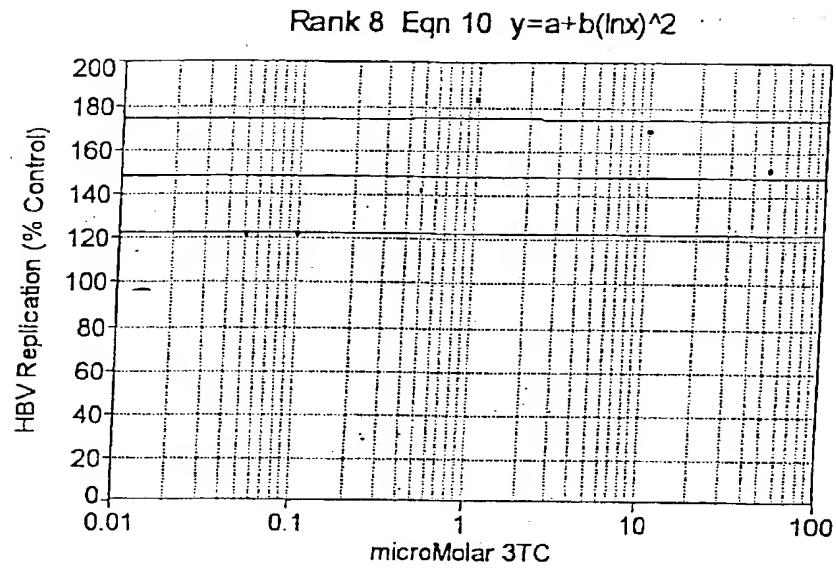


Figure 8A

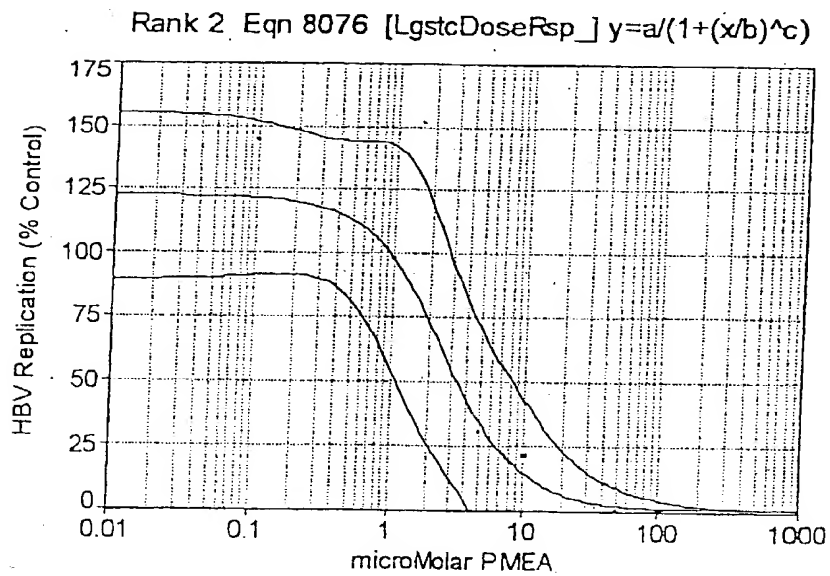


Figure 8B

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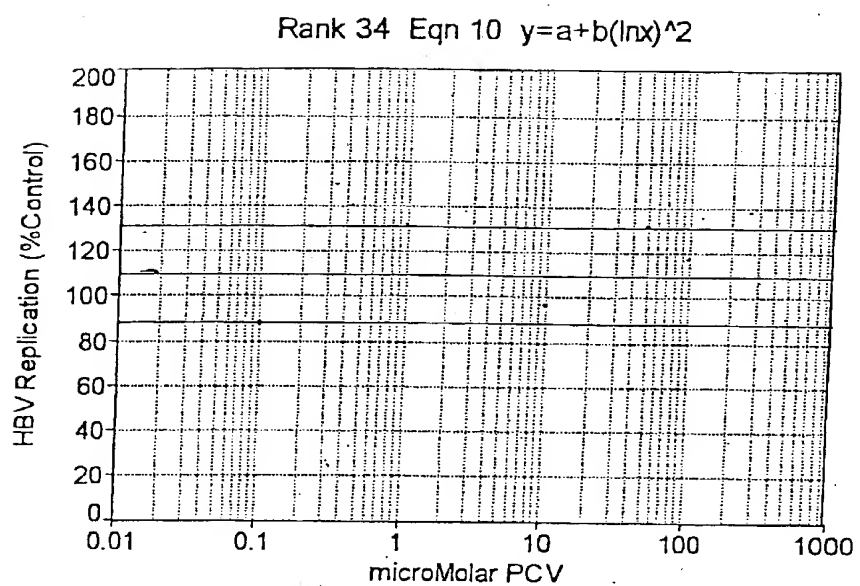


Figure 8C

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Rank 45 Eqn 10  $y=a+b(\ln x)^2$

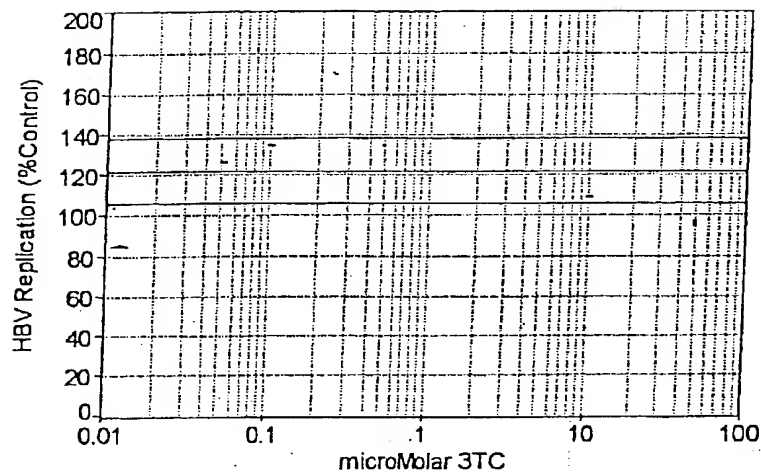


Figure 9A

Rank 2 Eqn 8076 [LgstcDoseRsp\_]  $y=a/(1+(x/b)^c)$

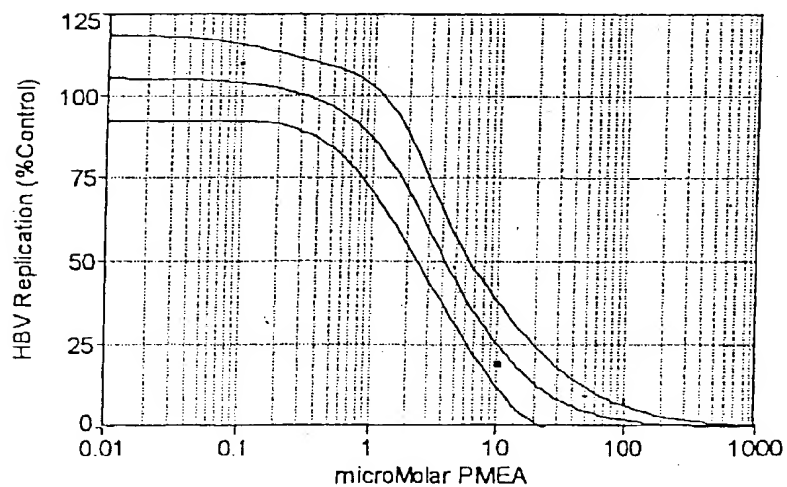


Figure 9B

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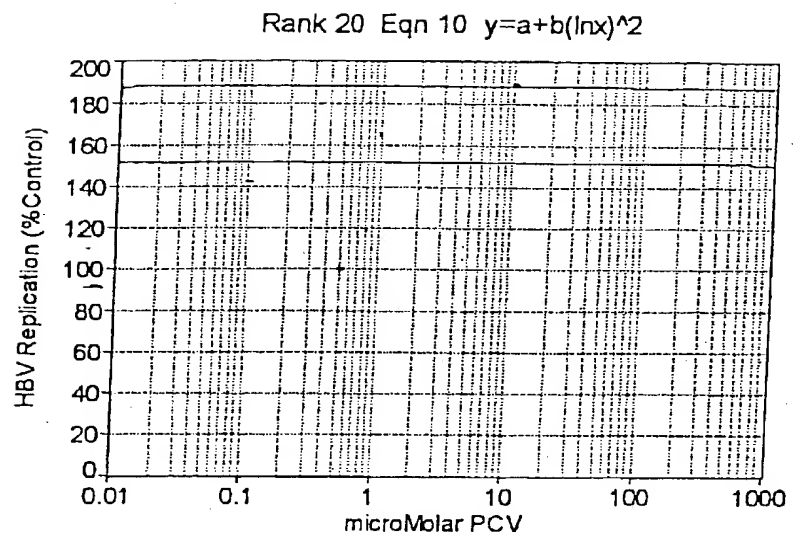


Figure 9C

## Cold dCTP Competition

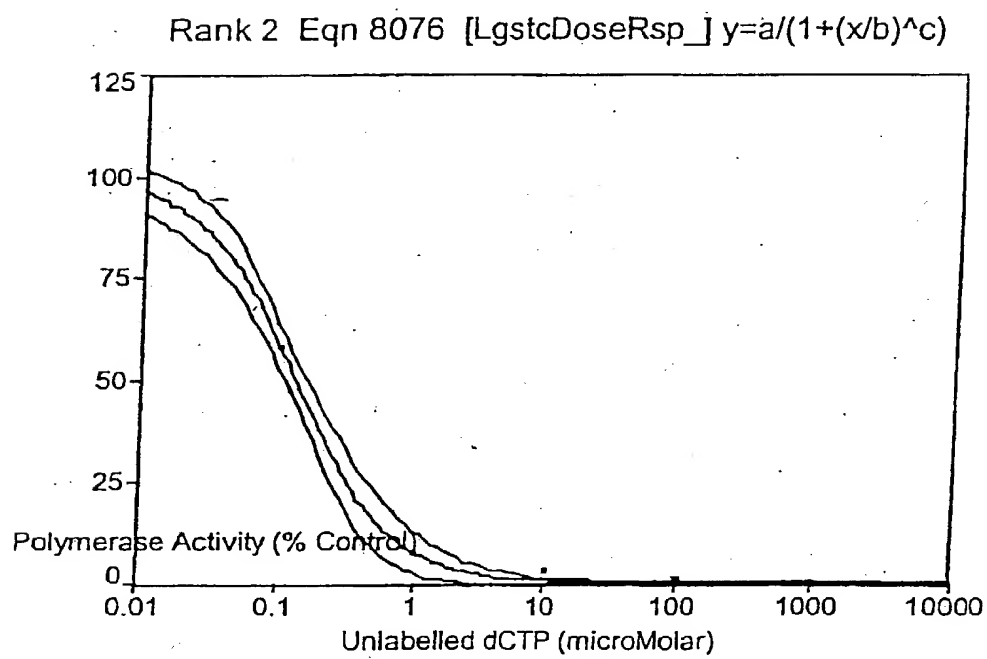


Figure 10



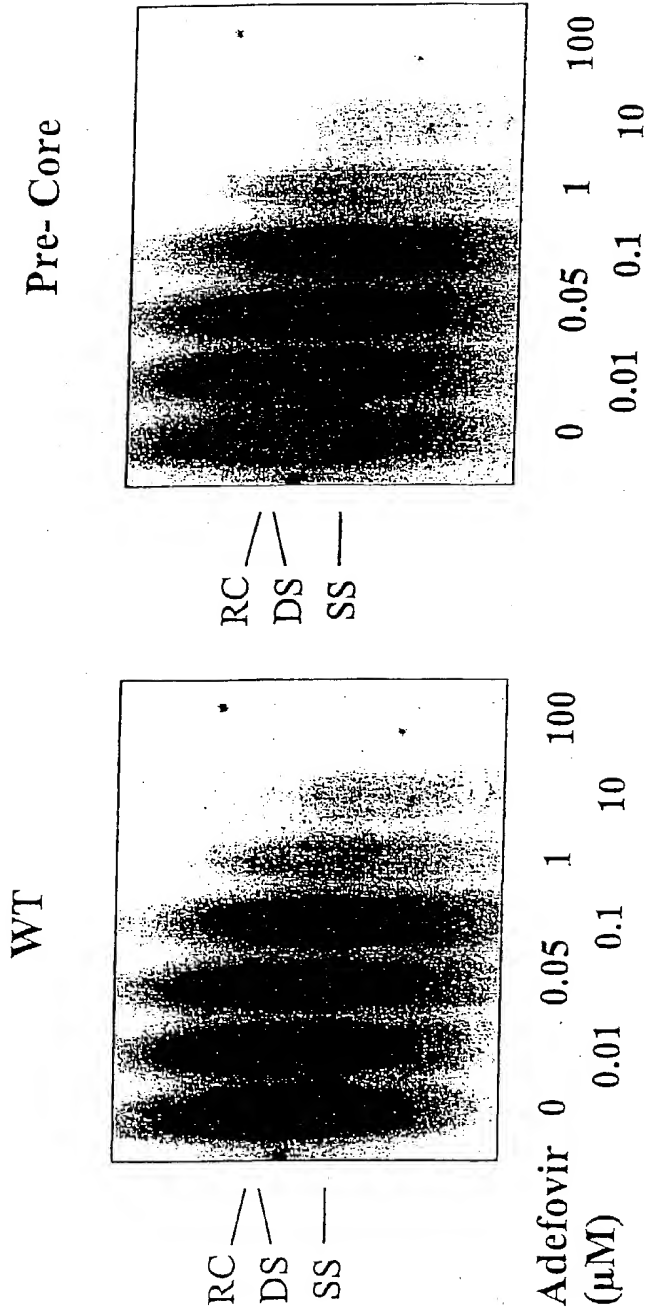


Figure 11A

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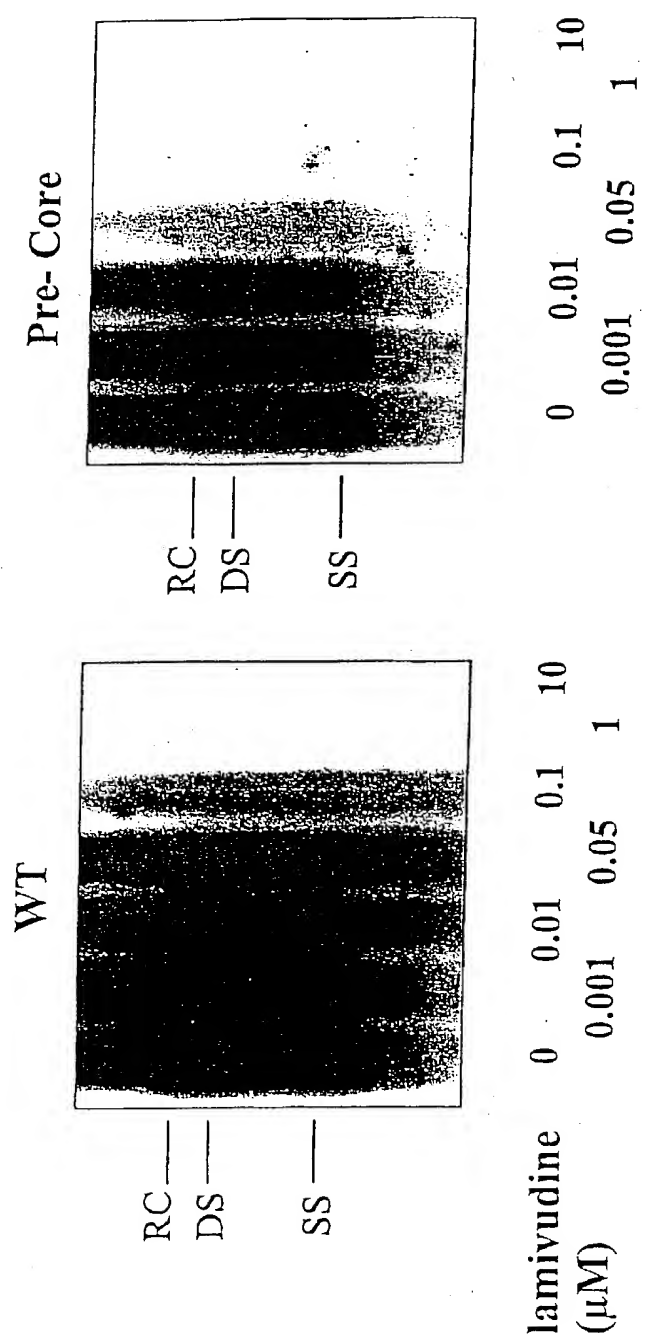


Figure 11B

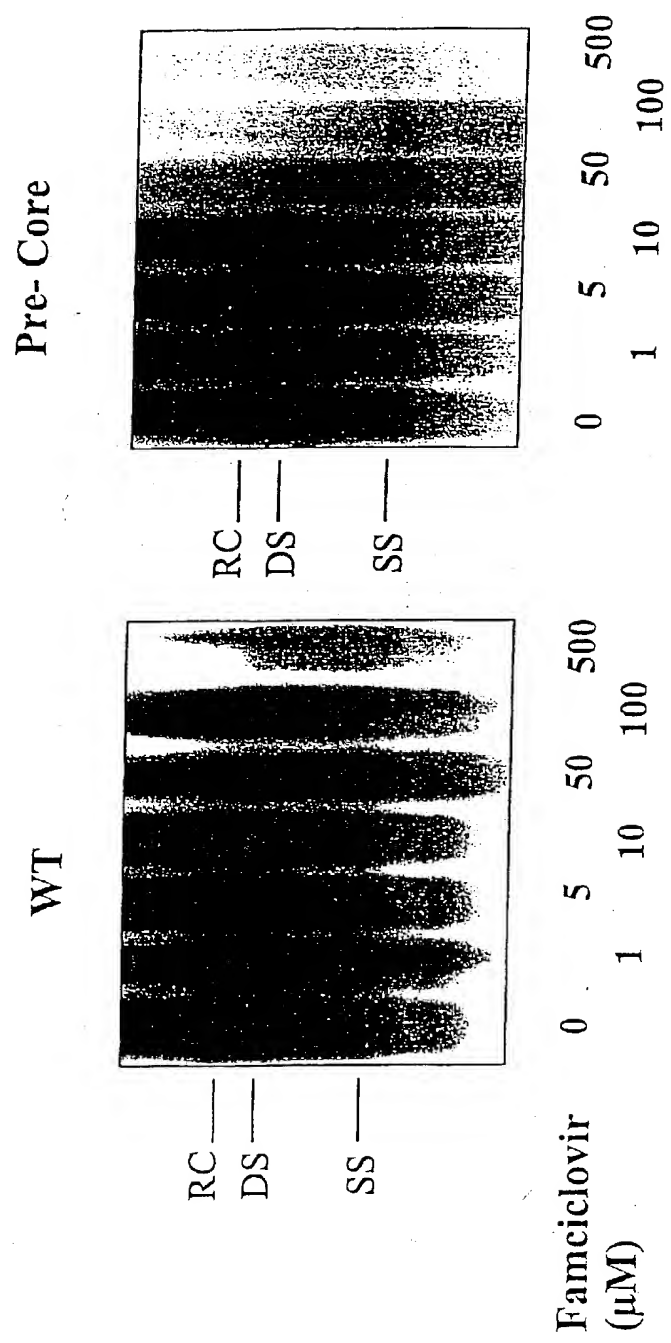


Figure 11C

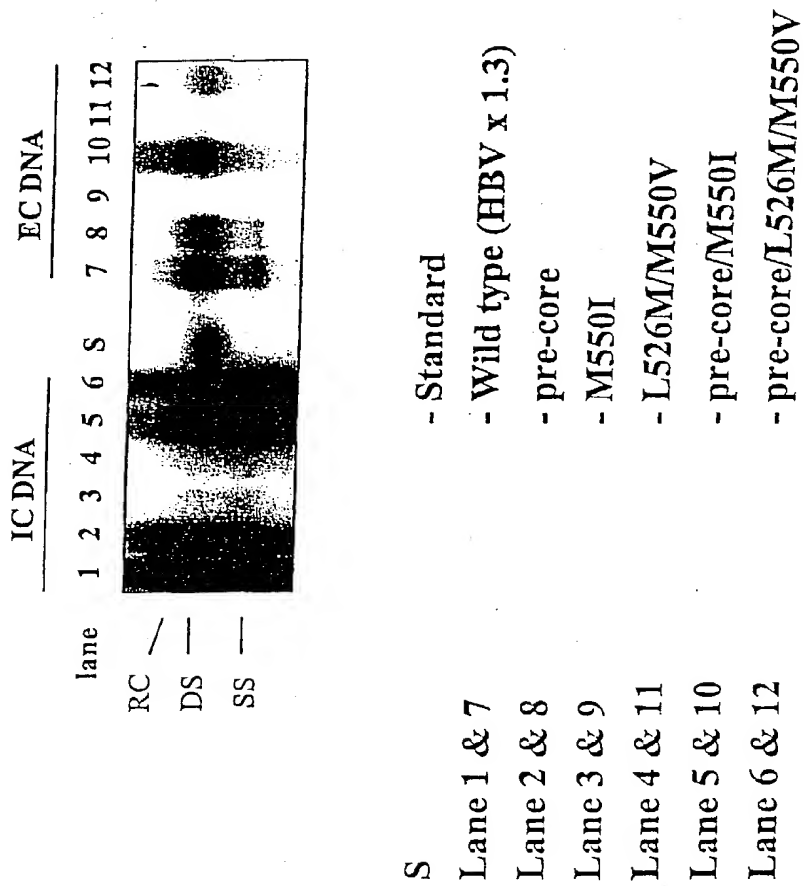


Figure 12

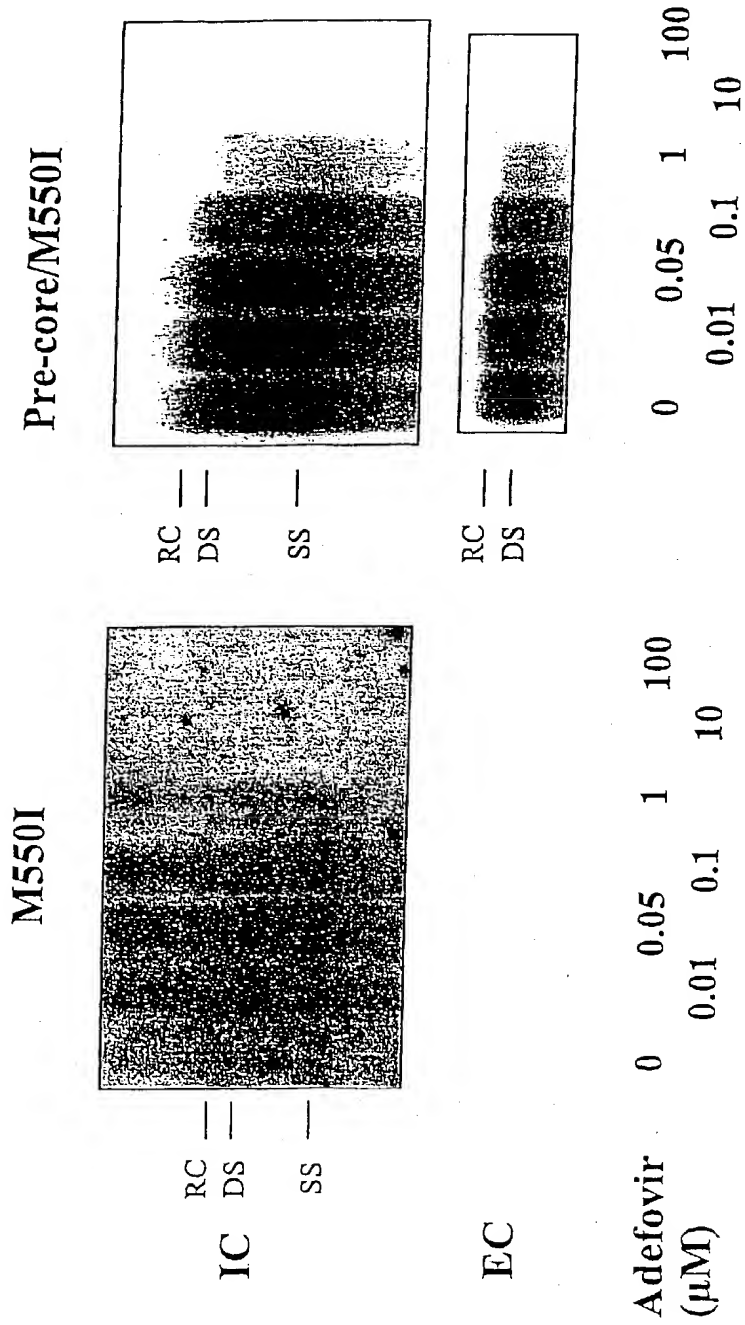


Figure 13A

pre-core/L526M/M550V

L526M/M550V

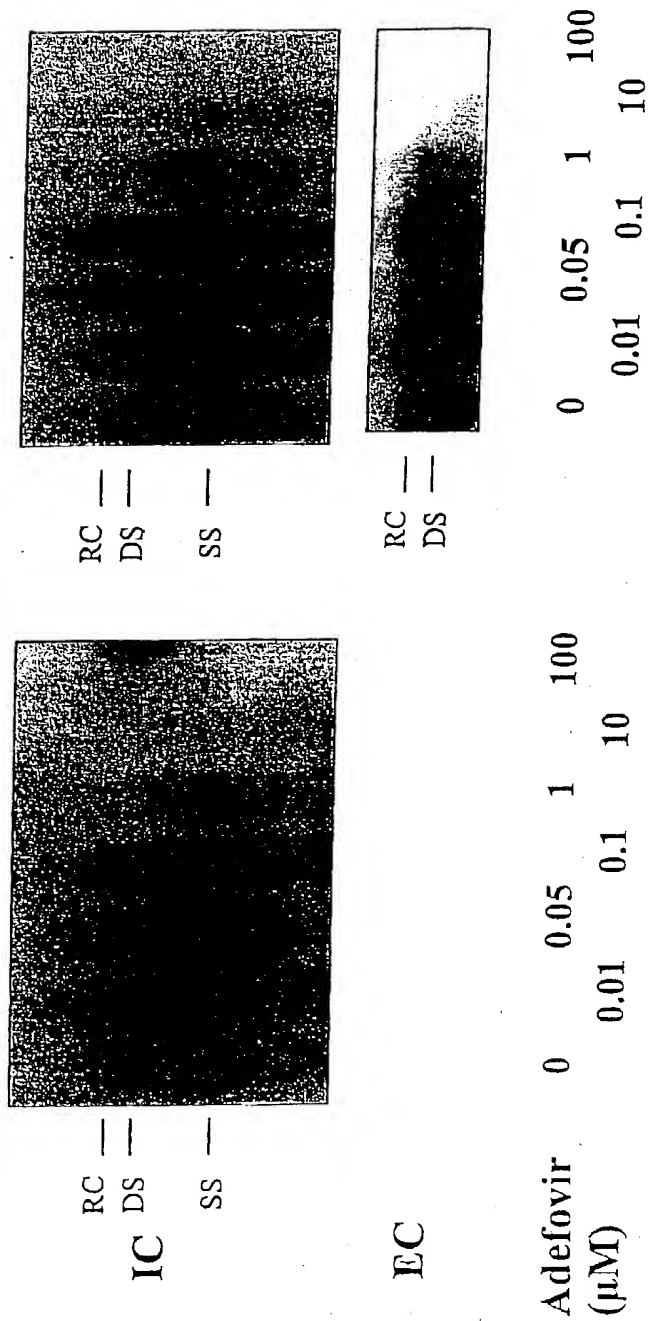
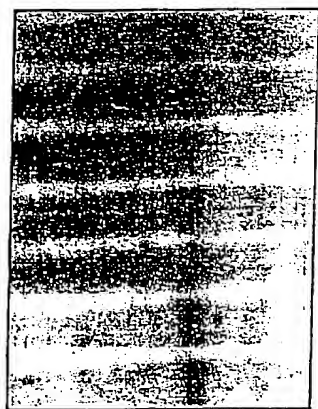


Figure 13B

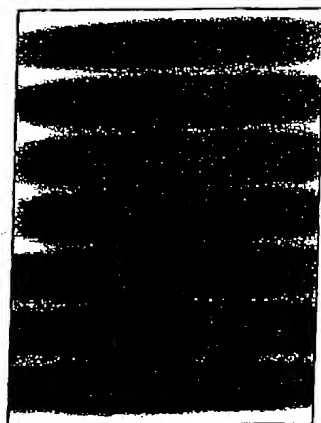
M550I



RC —  
DS —  
SS —

IC

Pre-core/M550I



RC —  
DS —  
SS —

EC



RC —  
DS —

lamivudine  
( $\mu$ M)

0 0.01 0.1 10  
0.001 0.05 1

0 0.01 0.1 10  
0.001 0.05 1

Figure 13C

L526M/M550V

Pre-core/L526M/M550V

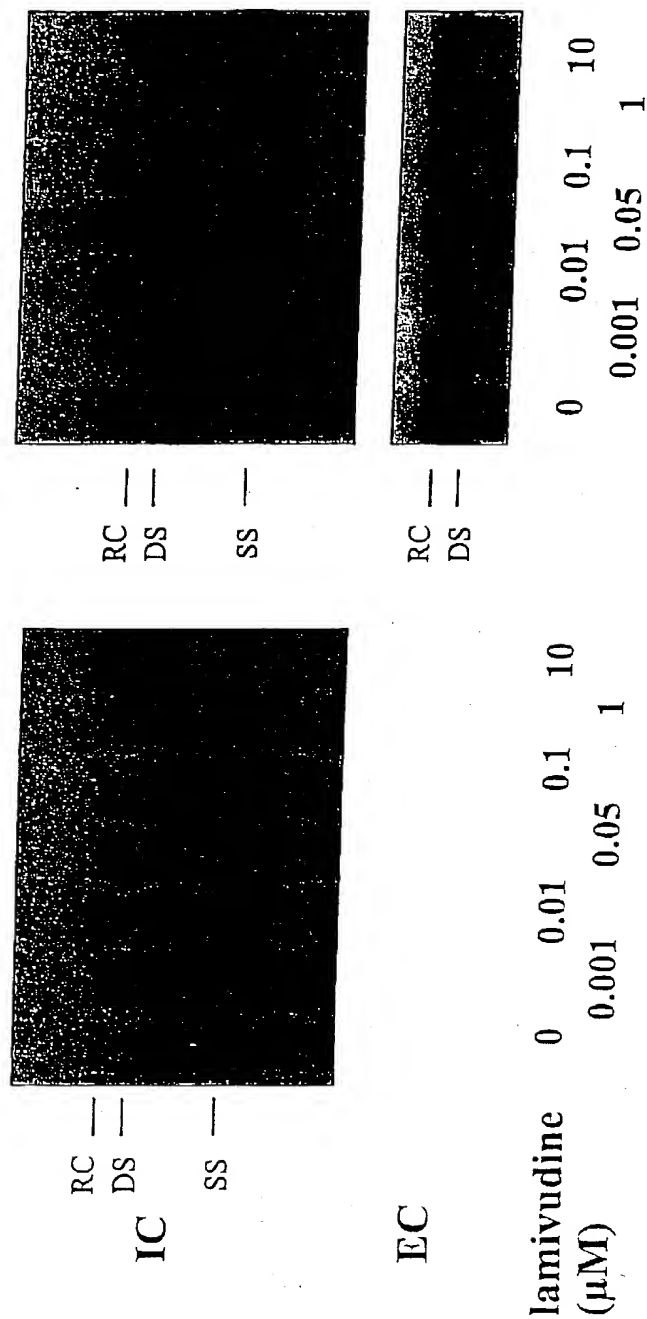
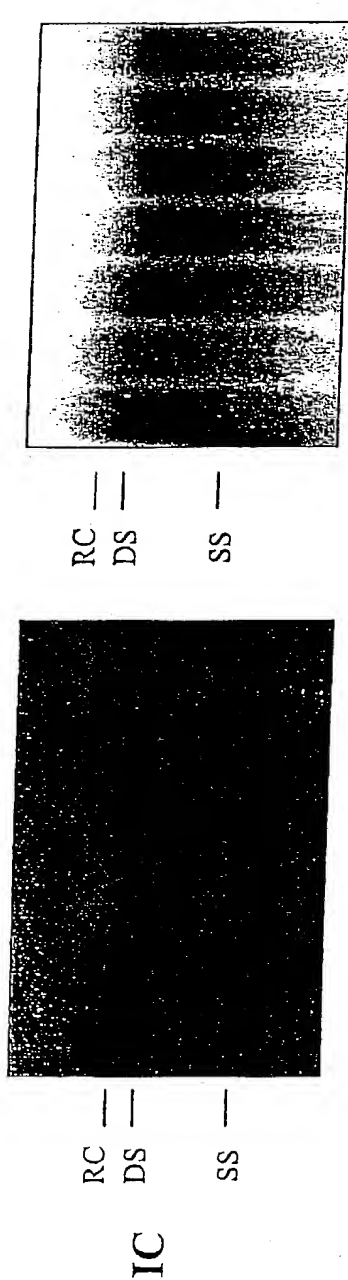


Figure 13D



M550I

Pre-core/M550I



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EC



Famciclovir  
( $\mu$ M)

0 5 50 500  
1 10 100 1000

Figure 13E

L526M/M550V      Pre-core/L526M/M550V

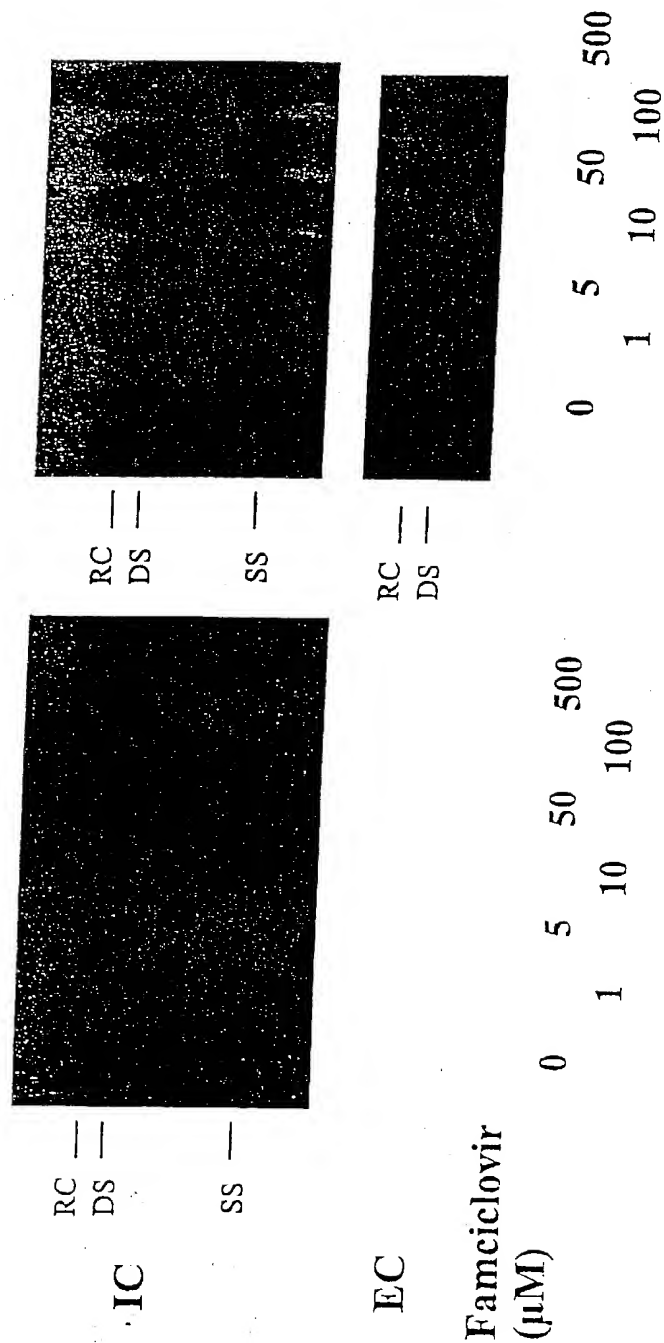


Figure 13F